0590 ##8 0107

OIPE

# 6

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/853,753

DATE: 12/17/2001 TIME: 15:04:49

Input Set : A:\ES.txt

Output Set: N:\CRF3\12172001\1853753.raw



3 <110> APPLICANT: Bech-Hansen, Torben 5 <120> TITLE OF INVENTION: GPI-Anchored Small Leucine-Rich Proteoglycan Gene NYX 7 <130> FILE REFERENCE: 45499-2 9 <140> CURRENT APPLICATION NUMBER: US 09/853,753 C--> 10 <141> CURRENT FILING DATE: 2001-05-17 12 <150> PRIOR APPLICATION NUMBER: CA 2,306,241 13 <151> PRIOR FILING DATE: 2000-05-12 15 <160> NUMBER OF SEQ ID NOS: 14 17 <170> SOFTWARE: PatentIn version 3.1 19 <210> SEO ID NO: 1 20 <211> LENGTH: 2297 21 <212> TYPE: DNA 22 <213> ORGANISM: Mus sp. 24 <300> PUBLICATION INFORMATION: 25 <301> AUTHORs: Bech-Hansen NT et al. 26 <302> TITLE: Mutations in NYX, encoding the leucine-rich proteoglycan nyctalopin, cause X-linked complete congenital stationary night blindness 28 <303> JOURNAL: Nature Genetics 29 <304> VOLUME: 26 30 <305> ISSUE: 3 31 <306> PAGES: 319-323 32 <307> DATE: 2000-11-01 33 <308> DATABASE ACCESSION NO: GenBank / AF254868 34 <309> DATABASE ENTRY DATE: 2000-12-23 36 <400> SEQUENCE: 1 37 ggctgaggga gtggaggggg acctcagagg agcaggacca gggagactcc caggacggta 60 39 ggggtcccac ggctgggtgg tcctaagcca ctgggtggat gaaaggccga gggatgttgg 120 41 teetgettet geatgeggtg gteeteggee tgeecagege etgggeegtg ggggeetgeg 180 43 cccgcgcttg tcccgccgcc tgcgcctgca gcaccgtgga gcgcggctgc tcggtgcgct 240 45 gegacegege gggeeteetg egggtgeegg eegageteee gtgegaggeg gteteeateg 300 47 acctggaccg gaacggcctg cgcttcctgg gcgagcgagc cttcggcacg ctgccgtcct 360 49 tgcgccgcct gtcgctgcgc cacaacaacc tgtccttcat cacgcccggc gccttcaagg 420 51 gcctgccgcg cctggctgag ctgcgcctgg cgcacaacgg cgacctgcgc tacctgcacg 480 53 cgcgcacctt cgcggcgctc agccgcctgc gccgcctaga cctagcagcc tgccgcctct 540 55 teagegtgee egagegeete etggeegaae tgeeggeeet gegegaaete geegeetteg 600 57 acaacctgtt ccgccgcgtg ccgggcgcc tgcgcggcct ggccaacctg acgcacgcgc 660 59 acctggagcg cggccgcatc gaggcggtgg cctccagctc gctgcagggc ctgcgccgc 720 61 tgcgctcgct cagcctgcag gccaaccgcg tccgtgccgt gcacgctggc gccttcgggg 780 63 actgtggcgt cctggagcat ctgctgctca acgacaacct gctggccgag ctcccggccg 840 65 acgcetteeg eggeetgegg egeetgegea egeteaacet gggtggeaac gegetggace 900 67 gegtggegeg egeetggtte getgaeetgg eegagetega getgetetae etggaeegea 960 69 acagcatege ettegtggag gagggegeet tecagaaeet etegggtete etegegetge 1020 71 accteaacgg caaccgcctc accgtgctcg cctgggtcgc cttccagccc ggcttcttcc 1080 73 tgggccgcct cttcctcttc cgcaacccgt ggtgctgcga ctgccgtctg gagtggctga 1140 75 gggactggat ggagggctcc ggacgtgtca ccgacgtgcc gtgcgcctcc ccgggctccg 1200 77 tggccggcct ggacctcagc caggtgacct tcgggcgctc ctccgatggc ctctgtgtgg 1260

79 accccgagga gctgaacctc accacgtcca gtccaggccc gtccccagaa ccagcggcca

1320

Input Set : A:\ES.txt

Output Set: N:\CRF3\12172001\1853753.raw

```
81 ccaccgtgag caggttcagc agcctcctct ccaagctgct ggccccgagg gtcccggtgg
83 aggaggcggc caacaccact ggggggctgg ccaacgcctc cctgtccgac agcctctcct
                                                                        1440
85 cccgtggggt gggaggcgcg ggccggcagc cctggtttct cctcqcctct tqtctcctqc
                                                                        1500
87 ccagcgtggc ccagcacgtg gtgtttggcc tgcagatgga ctgacctggc cagagggggg
                                                                        1560
89 aaagtttgct taactgggct tgagtgtgtt tgtggtaagg ggagaggagc cggaatggag
                                                                        1620
91 ggcagaggtg aaaatcccag tggagggtgg aaggaaccgt ttgcctccag agatggcccc
                                                                        1680
93 agggagaaca cagggacgtg ccactcgagg gggaggatgg tatggatttc tgcttttgtc
                                                                        1740
95 acacgggcat ccattggaaa agagaagcaa gaatgaacgt gggccctcgg gtgggaagac
                                                                        1800
97 taggaatcgg aagcttctag ggcttcacat cccttcccct ccctcccct tcccctcatc
                                                                        1860
99 ttccaggcaa cagtgcctgc aaggcctgaa ttagagagac ttccattggc taagtagtta
101 agagccgtcc catttctcct ggcggggtaa cccattacac cgaagtcctt tgttttctac
                                                                         1980
103 cacaatcctc ctcctctt ccaggggcct ggaaacacta ggattcagga aggtaggcag
                                                                         2040
105 gacgtgagag aagggagatg ggagagagat ttaagacaaa gggtggcggt ggttcctggg
                                                                         2100
107 gtctgagatg tgttaggagg cgtttaaaac aaagatccag ttcatttact ccacagttat
                                                                         2160
109 tcccagggct ggccctagcc acaaaggaac tttagggcag ggtagggaaa aaaggggcag
                                                                         2220
111 cagggggtgt gtttgtggac aaataaattt gtaaagtccg aggattaaaa aaaaaaaaa
                                                                         2280
113 gttaaaccgg tttctct
                                                                         2297
116 <210> SEQ ID NO: 2
117 <211> LENGTH: 481
118 <212> TYPE: PRT
119 <213> ORGANISM: Mus sp.
121 <400> SEQUENCE: 2
123 Met Lys Gly Arg Gly Met Leu Val Leu Leu His Ala Val Val Leu
124 1
127 Gly Leu Pro Ser Ala Trp Ala Val Gly Ala Cys Ala Arg Ala Cys Pro
128
                20
                                    25
131 Ala Ala Cys Ala Cys Ser Thr Val Glu Arg Gly Cys Ser Val Arg Cys
                                40
135 Asp Arg Ala Gly Leu Leu Arg Val Pro Ala Glu Leu Pro Cys Glu Ala
136
                            55
139 Val Ser Ile Asp Leu Asp Arg Asn Gly Leu Arg Phe Leu Gly Glu Arg
140 65
                        70
                                            75
143 Ala Phe Gly Thr Leu Pro Ser Leu Arg Arg Leu Ser Leu Arg His Asn
                                        90
147 Asn Leu Ser Phe Ile Thr Pro Gly Ala Phe Lys Gly Leu Pro Arg Leu
148
                100
                                    105 ·
151 Ala Glu Leu Arg Leu Ala His Asn Gly Asp Leu Arg Tyr Leu His Ala
152
                                120
                                                    125
155 Arg Thr Phe Ala Ala Leu Ser Arg Leu Arg Arg Leu Asp Leu Ala Ala
156
       130
                            135
159 Cys Arg Leu Phe Ser Val Pro Glu Arg Leu Leu Ala Glu Leu Pro Ala
                        150
                                            155
163 Leu Arg Glu Leu Ala Ala Phe Asp Asn Leu Phe Arg Arg Val Pro Gly
164
                    165
                                        170
167 Ala Leu Arg Gly Leu Ala Asn Leu Thr His Ala His Leu Glu Arg Gly.
168
                180
                                    185
171 Arg Ile Glu Ala Val Ala Ser Ser Leu Gln Gly Leu Arg Arg Leu
                                200
175 Arg Ser Leu Ser Leu Gln Ala Asn Arg Val Arg Ala Val His Ala Gly
```

Input Set : A:\ES.txt

Output Set: N:\CRF3\12172001\1853753.raw

```
176
        210
179 Ala Phe Gly Asp Cys Gly Val Leu Glu His Leu Leu Leu Asn Asp Asn
                        230
                                            235
183 Leu Leu Ala Glu Leu Pro Ala Asp Ala Phe Arg Gly Leu Arg Arg Leu
                   245
                                        250
187 Arg Thr Leu Asn Leu Gly Gly Asn Ala Leu Asp Arg Val Ala Arg Ala
                260
                                    265
191 Trp Phe Ala Asp Leu Ala Glu Leu Glu Leu Tyr Leu Asp Arg Asn
                                280
195 Ser Ile Ala Phe Val Glu Glu Gly Ala Phe Gln Asn Leu Ser Gly Leu
199 Leu Ala Leu His Leu Asn Gly Asn Arg Leu Thr Val Leu Ala Trp Val
                        310
                                            315
203 Ala Phe Gln Pro Gly Phe Phe Leu Gly Arg Leu Phe Leu Phe Arg Asn
                    325
                                        330
207 Pro Trp Cys Cys Asp Cys Arg Leu Glu Trp Leu Arg Asp Trp Met Glu
                340
                                    345
211 Gly Ser Gly Arg Val Thr Asp Val Pro Cys Ala Ser Pro Gly Ser Val
                                360
                                                    365
215 Ala Gly Leu Asp Leu Ser Gln Val Thr Phe Gly Arg Ser Ser Asp Gly
       370
                            375
219 Leu Cys Val Asp Pro Glu Glu Leu Asn Leu Thr Thr Ser Ser Pro Gly
                        390
                                            395
223 Pro Ser Pro Glu Pro Ala Ala Thr Thr Val Ser Arg Phe Ser Ser Leu
                    405
                                        410
227 Leu Ser Lys Leu Leu Ala Pro Arg Val Pro Val Glu Glu Ala Ala Asn
228
                420
                                    425
231 Thr Thr Gly Gly Leu Ala Asn Ala Ser Leu Ser Asp Ser Leu Ser Ser
           435
                                440
235 Arg Gly Val Gly Gly Ala Gly Arg Gln Pro Trp Phe Leu Leu Ala Ser
236 450
                            455
239 Cys Leu Leu Pro Ser Val Ala Gln His Val Val Phe Gly Leu Gln Met
240 465
                        470
                                            475
243 Asp
247 <210> SEQ ID NO: 3
248 <211> LENGTH: 20
249 <212> TYPE: DNA
250 <213> ORGANISM: Artificial Sequence
252 <220> FEATURE:
253 <223> OTHER INFORMATION: PCR primer
255 <220> FEATURE:
256 <221> NAME/KEY: misc_feature
257 <222> LOCATION: (1)..(20)
258 <223> OTHER INFORMATION: forward primer for polymorphism 506B13CA1 (DXS10042)
261 <400> SEQUENCE: 3
262 atcacagtgc cctgcctaaa
                                                                           20
265 <210> SEQ ID NO: 4
266 <211> LENGTH: 20
267 <212> TYPE: DNA
```

Input Set : A:\ES.txt

Output Set: N:\CRF3\12172001\1853753.raw

268 <213> ORGANISM: Artificial Sequence 270 <220> FEATURE: 271 <223> OTHER INFORMATION: PCR primer 273 <220> FEATURE: 274 <221> NAME/KEY: misc\_feature 275 <222> LOCATION: (1)..(20) 276 <223> OTHER INFORMATION: reverse primer for polymorphism 506B13CA (DXS10042) 279 <400> SEQUENCE: 4 280 tcccaaagtg ctgggattac 20 283 <210> SEQ ID NO: 5 284 <211> LENGTH: 21 285 <212> TYPE: DNA 286 <213> ORGANISM: Artificial Sequence 288 <220> FEATURE: 289 <223> OTHER INFORMATION: PCR primer 291 <220> FEATURE: 292 <221> NAME/KEY: misc\_feature 293 <222> LOCATION: (1)..(21) 294 <223> OTHER INFORMATION: forward primer for polymorphism 200L4CA1 (DXS10044) 297 <400> SEQUENCE: 5 298 gaacagcaaa ccaaatccaa a 21 301 <210> SEQ ID NO: 6 302 <211> LENGTH: 20 303 <212> TYPE: DNA 304 <213> ORGANISM: Artificial Sequence 306 <220> FEATURE: 307 <223> OTHER INFORMATION: PCR primer 309 <220> FEATURE: 310 <221> NAME/KEY: misc\_feature 311 <222> LOCATION: (1)..(20) 312 <223> OTHER INFORMATION: reverse primer for polymorphism 200L4CA1 (DXS10044) 315 <400> SEQUENCE: 6 316 ggcctatggt aatgcctcct 20 319 <210> SEQ ID NO: 7 320 <211> LENGTH: 20 321 <212> TYPE: DNA 322 <213> ORGANISM: Artificial Sequence 324 <220> FEATURE: 325 <223> OTHER INFORMATION: PCR primer 327 <220> FEATURE: 328 <221> NAME/KEY: misc\_feature 329 <222> LOCATION: (1)..(20) 330 <223> OTHER INFORMATION: forward primer for polymorphism 169I5CA2 (DXS10045) 333 <400> SEQUENCE: 7 334 aaacttagct gggcatgctg 20 337 <210> SEQ ID NO: 8 338 <211> LENGTH: 21

339 <212> TYPE: DNA

340 <213> ORGANISM: Artificial Sequence

Input Set : A:\ES.txt

Output Set: N:\CRF3\12172001\1853753.raw

342 <220> FEATURE: 343 <223> OTHER INFORMATION: PCR primer 345 <220> FEATURE: 346 <221> NAME/KEY: misc\_feature 347 <222> LOCATION: (1)..(21) 348 <223> OTHER INFORMATION: reverse primer for polymorphism 169I5CA2 (DXS10045) 351 <400> SEQUENCE: 8 352 gctgggacta catacagcac a 21 355 <210> SEQ ID NO: 9 356 <211> LENGTH: 21 357 <212> TYPE: DNA 358 <213> ORGANISM: Artificial Sequence 360 <220> FEATURE: 361 <223> OTHER INFORMATION: PCR primer 363 <220> FEATURE: 364 <221> NAME/KEY: misc\_feature 365 <222> LOCATION: (1)..(21) 366 <223> OTHER INFORMATION: forward primer for NYX expression 369 <400> SEQUENCE: 9 370 agggagtgga ggggacctca g 21 373 <210> SEQ ID NO: 10 374 <211> LENGTH: 18 375 <212> TYPE: DNA 376 <213> ORGANISM: Artificial Sequence 378 <220> FEATURE: 379 <223> OTHER INFORMATION: PCR primer 381 <220> FEATURE: 382 <221> NAME/KEY: misc\_feature 383 <222> LOCATION: (1)..(18) 384 <223> OTHER INFORMATION: reverse primer for NYX expression 387 <400> SEQUENCE: 10 388 acqqcacqqa cqcqqttq 18 391 <210> SEQ ID NO: 11 392 <211> LENGTH: 20 393 <212> TYPE: DNA 394 <213> ORGANISM: Artificial Sequence 396 <220> FEATURE: 397 <223> OTHER INFORMATION: PCR primer 399 <220> FEATURE: 400 <221> NAME/KEY: misc\_feature 401 <222> LOCATION: (1)..(20) 402 <223> OTHER INFORMATION: forward primer for W350X mutation 405 <400> SEQUENCE: 11 406 gatttttcct ggggtgacct 20 409 <210> SEQ ID NO: 12 410 <211> LENGTH: 19 411 <212> TYPE: DNA 412 <213> ORGANISM: Artificial Sequence

414 <220> FEATURE:

VERIFICATION SUMMARY

DATE: 12/17/2001

PATENT APPLICATION: US/09/853,753

TIME: 15:04:50

Input Set : A:\ES.txt

Output Set: N:\CRF3\12172001\1853753.raw

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date